

<110> DeAngelis, Paul Jing, Wei

<120> TARGETED GLYCOSAMINOGLYCAN POLYMERS BY POLYMER GRAFTING AND METHODS OF MAKING AND USING SAME

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<150> 60/404,356

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Ile Ile Met Thr Ser His Asn Thr Glu Lys Phe Ile Glu Ala Ser Ile 100 105 110

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Val Asn Ala Leu Leu Ser Asn Lys Asp Asn Ile Ala Val Arg Cys Ala Page 23

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205

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Pro Ala Asp Tyr Thr Asn Thr Met Ile Lys Lys Ile Asn Lys Tyr Asn

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Ile Gln Ser Lys Leu Ile Ile Ser Asn Asn Pro Trp Gly Tyr Ser Ser Page 25

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<213> Pasteurella multocida

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Page 41

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1800

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1260

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1860

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Page 52

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Page 55

1860

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1860

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1200

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<223> synthetic peptide based on residues 526-543 of pmHAS

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Lys Glu

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<211> 19

<212> PRT

<213> artificial sequence

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<213> Pasteurella multocida

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1380

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Page 74

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<211> 2007

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1	\sim	\sim	^
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Page 78

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Page 81

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Page 84

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1620

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1620

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Page 122

1200

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1680

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Maria Salahan

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Gln Ser Lys Ser Asn Lys Ile Glu Glu Asp Asn Ile Ser Gly Glu Asn 50 55 60

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Glu Leu Gly Ile Thr Lys Glu Arg Leu Gly Ala Pro Pro Leu Val Ser 85 90 95

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415

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900

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 105 110

Arg Glu Val Ser Ser Gly Ser Met Glu Gly Leu Trp Asn Thr Phe Thr 195 200 205

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Ile His Lys Arg Ile Asn Glu Tyr Asp Asn Val Leu Glu Leu Ser Lys 50 55 60

Asn Val Ser Ala Gln Asn Ser Gly Asn Glu Phe Ser Tyr Leu Leu Gly 65 70 75 80

Tyr Ala Asp Ser Leu Arg Lys Val Gly Met Leu Asp Thr Tyr Ile Lys 85 90 95

Ile Val	Cys	Tyr	Leu	Thr	Ile	Gln	Ser	Arg	Tyr	Phe	Lys	Asn	Gly	Glu
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Arg	Val	Lys	Leu	Phe	Glu	His	Ile	Ser	Asn	Ala	Leu	Arg	Tyr	Ser	Arg
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Asn	Ala	Asn	Asp	Asp	Met	Gln	Asp	Ser	Tyr	Asn	Leu	Leu	Pro	Glu	Gln
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Ser	Gln	Phe	Asn	Lys	Cys	Leu	Arg	Lys	Tyr	Asp	Leu	Ser	Glu	Ile	Thr
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Phe Asn Ser Glu Asp Thr Ile Ala Tyr Ser Leu His Ser Leu Leu Asn 275 280 285

SEQUENCE LISTING.ST25.txt Gln Thr Tyr Glu Asn Ile Glu Ile Leu Val Cys Asp Asp Cys Ser Ser 290 295 300

Asp Lys Ser Leu Glu Ile Ile Lys Ser Ile Ala Tyr Ser Ser Ser Arg 305 310 315 320

Val Lys Val Tyr Ser Ser Arg Lys Asn Gln Gly Pro Tyr Asn Ile Arg 325 330 335

Asn Glu Leu Ile Lys Lys Ala His Gly Asn Phe Ile Thr Phe Gln Asp 340 345 350

Ala Asp Asp Leu Ser His Pro Glu Arg Ile Gln Arg Gln Val Glu Val 355 360 365

Leu Arg Asn Asn Lys Ala Val Ile Cys Met Ala Asn Trp Ile Arg Val 370 375 380

Ala Ser Asn Gly Lys Ile Gln Phe Phe Tyr Asp Asp Lys Ala Thr Arg 385 390 395 400

Met Ser Val Val Ser Ser Met Ile Lys Lys Asp Ile Phe Ala Thr Val 405 410 415

Gly Gly Tyr Arg Gln Ser Leu Ile Gly Ala Asp Thr Glu Phe Tyr Glu 420 425 430

Thr Val Ile Met Arg Tyr Gly Arg Glu Ser Ile Val Arg Leu Leu Gln
435 440 445

Pro Leu Ile Leu Gly Leu Trp Gly Asp Ser Gly Leu Thr Arg Asn Lys 450 455 460

Gly Thr Glu Ala Leu Pro Asp Gly Tyr Ile Ser Gln Ser Arg Arg Glu 465 470 475 480

Tyr Ser Asp Ile Ala Ala Arg Gln Arg Val Leu Gly Lys Ser Ile Val 485 490 495

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<212> PRT

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Ser His Ser Arg Arg Glu Glu His Ser Gly Arg Asn Gly Leu His Gln
35 40 45

Pro Ser Pro Asp His Phe Trp Pro Arg Phe Pro Asp Ala Leu Arg Pro 50 55 60

Phe Phe Pro Trp Asp Gln Leu Glu Asn Glu Asp Ser Ser Val His Ile 70 75 80

Ser Pro Arg Gln Lys Arg Asp Ala Asn Ser Ser Ile Tyr Lys Gly Lys 85 90 95

Lys Cys Arg	Met Glu	Ser	Cys	Phe	Asp	Phe	Thr	Leu	Cys	Lys	Lys	Asn
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Gly	Phe	Lys	Val	Tyr	Val	Tyr	Pro	Gln	Gln	Lys	Gly	Glu	Lys	Ile	Ala
		115					120					125			

Phe	Asp	Ile	Gly	Gln	Ala	Met	Leu	Ala	Lys	Ala	Ser	Ile	Ser	Thr	Glu
	210					215			_		220				

Asn	Phe	Arg	Pro	Asn	Phe	Asp	Val	Ser	Ile	Pro	Leu	Phe	Ser	Lys	Asp
225					230					235					240

Thr Gly Ile Gly Ser Asp Thr Arg Asn Ala Leu Tyr His Val His Asn 275 280 285

Gly	Glu	Asp	Val	Leu	Leu	Leu	Thr	Thr	Cys	Lys	His	Gly	Lys	Asp	Trp
_	290					295	•				300				•

Gln	Lys	His	Lys	Asp	Ser	Arg	Cys	Asp	Arg	Asp	Asn	Thr	Glu	Tyr	Glu
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Thr Ala Val Ile His Ala Val Thr Pro Leu Val Ser Gln Ser Gln Pro

				485					490					495	
Val	Leu	Lys	Leu 500	Leu	Val	Ala	Ala	Ala 505	Lys	Ser	Gln	Tyr	Cys 510	Ala	Gln
Ile	Ile	Val 515	Leu	Trp	Asn	Cys	Asp 520	Lys	Pro	Leu	Pro	Ala 525	Lys	His	Arg
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Val 545	Met	Ser	Ser	Arg	Phe 550	Leu	Pro	Tyr	Asp	Asn 555	Ile	Ile	Thr	Asp	Ala 560
Val	Leu	Ser	Leu	Asp 565	Glu	Asp	Thr	Val	Leu 570	Ser	Thr	Thr	Glu	Val 575	Asp
Phe	Ala	Phe	Thr 580	Val	Trp	Gln	Ser	Phe 585	Pro	Glu	Arg	Ile	Val 590	Gly	Tyr

Pro Ala Arg Ser His Phe Trp Asp Asn Ser Lys Glu Arg Trp Gly Tyr 595 600 605

Thr Ser Lys Trp Thr Asn Asp Tyr Ser Met Val Leu Thr Gly Ala Ala 610 620

Ile Tyr His Lys Tyr Tyr His Tyr Leu Tyr Ser His Tyr Leu Pro Ala 625 630 635 640

Ser Leu Lys Asn Met Val Asp Gln Leu Ala Asn Cys Glu Asp Ile Leu 645 650 655

Met Asn Phe Leu Val Ser Ala Val Thr Lys Leu Pro Pro Ile Lys Val 660 665 670

Thr Gln Lys Lys Gln Tyr Lys Glu Thr Met Met Gly Gln Thr Ser Arg
675 680 685

Ala Ser Arg Trp Ala Asp Pro Asp His Phe Ala Gln Arg Gln Ser Cys 690 695 700

Met Asn Thr Phe Ala Ser Trp Phe Gly Tyr Met Pro Leu Ile His Ser 705 710 715 720

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Arg Lys Lys Tyr Arg Asp Ile Glu Arg Leu 740 745

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<212> PRT

<213> Mus musculus

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Arg Met Lys Thr Lys His Arg Ile Tyr Tyr Val Thr Leu Phe Ser Ile 20 25 30

Val Leu Gly Leu Ile Ala Thr Gly Met Phe Gln Phe Trp Pro His 35 40 45

Ser Ile Glu Ser Ser Ser Asp Gly Gly Val Glu Lys Arg Ser Ile Arg 50 55 60

SEQUENCE LISTING.ST25.txt Glu Val Pro Val Val Arg Leu Pro Thr Asp Ser Pro Ile Pro Glu Arg 65 70 75 80 Gly Asp Leu Ser Cys Arg Met His Thr Cys Phe Asp Val Tyr Arg Cys

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95

Gly	Phe	Asn	Pro	Lys	Asn	Lys	Ile	Lys	Val	Tyr	Ile	Tyr	Pro	Leu	Lys
			100					105			-		110		

85

Lys Tyr Val Asp Asp Ala Gly Val Pro Val Ser Ser Ala Ile Ser Arg 115 120 125

Glu Tyr Asn Glu Leu Leu Thr Ala Ile Ser Asp Ser Asp Tyr Tyr Thr 130 135 140

Asp Asp Ile Asn Arg Ala Cys Leu Phe Val Pro Ser Ile Asp Val Leu 145 150 155 160

Asn Gln Asn Pro Leu Arg Ile Lys Glu Thr Ala Gln Ala Leu Ala Gln 165 170 175

Leu Ser Arg Trp Asp Arg Gly Thr Asn His Leu Leu Phe Asn Met Leu 180 185 190

Pro Gly Ala Pro Pro Asp Tyr Asn Thr Ala Leu Asp Val Pro Arg Asp 195 200 205

Arg Ala Leu Leu Ala Gly Gly Gly Phe Ser Thr Trp Thr Tyr Arg Gln 210 215 220

Gly Tyr Asp Val Ser Ile Pro Val Phe Ser Pro Leu Ser Ala Glu Met 225 230 235 240

Ala Leu Pro Glu Lys Ala Pro Gly Pro Arg Arg Tyr Phe Leu Leu Ser 245 250 255

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Gln	Ala	Lys	His	Gln	Glu	Ser	Val	Leu	Val	Leu	Asp	Lys	Cys	Thr	Asn
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Leu Arg Arg Ala Arg Leu Gly Gln Ala Val Leu Ser Asp Val Leu Gln
$$325$$
 330 335

Lys	Met	Ser	Asp	Val	Tyr	Ser	Ile	Leu	Gln	Asn	Ile	Pro	Gln	Arg	Gln
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Pro Ala Val Lys Trp Ala Ser Val Ser Asn Pro Leu Phe Leu Pro Leu 435 440 445

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Arg	Val	Glu	Ser	Leu	Phe	Arg	Val	Ile	Thr	Glu	Val	Ser	Lys	Val	Pro
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Cys Thr Ala Ile Asp Gly Leu Ser Leu Asp Gln Thr His Met Val Glu 660 665 670

Arg Ser Glu Cys Ile Asn Lys Phe Ala Ser Val Phe Gly Thr Met Pro 675 680 685

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Asp Xaa Xaa Xaa Xaa Xaa Xaa Xaa Ile Ala Xaa Xaa Xaa Xaa Xaa 20 25 30

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acid
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Page 163

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1680

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<213> Pasteurella multocida

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Gln Ser Lys Ser Asn Lys Ile Glu Glu Asp Asn Ile Ser Gly Glu Asn 50 55 60

Lys Phe Ser Val Ser Ile Lys Asp Leu Tyr Asn Glu Ile Ser Asn Ser 65 70 75 80

Glu Leu Gly Ile Thr Lys Glu Arg Leu Gly Ala Pro Pro Leu Val Ser 85 90 95

Ile Ile Met	Thr	Ser	His	Asn	Thr	Glu	Lys	Phe	Ile	Glu	Ala	Ser	Ile
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Asn	Ser	Leu	Leu	Leu	Gln	Thr	Tyr	Asn	Asn	Leu	Glu	Val	Ile	Val	Val
		115					120					125			

Cys	Ala	Tyr	Ser	Arg	Ile	Asn	Leu	Glu	Thr	Gln	Asn	Ile	Ile	Lys	Val
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Asn	Asp	Asn	Lys	Tyr	Lys	Leu	Gly	Leu	Ile	Thr	Leu	Gly	Val	Tyr	Arg
225					230					235					240

Ile Asn Asn Leu Phe Leu Pro Leu Tyr Tyr Asn Thr Met Arg Glu Asp 275 280 285

Ser	Leu	Phe	Ser	Asp	Met	Val	Glu	Trp	Val	Asp	Glu	Asn	Asn	Ile	Lys
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Gln	Lys	Thr	Ser	Asp	Ala	Arg	Gln	Asn	Tyr	Leu	His	Glu	Phe	Gln	Lys
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Lys Pro Leu Glu Asn Asp Thr Ala Val Asn Ile Leu Gly Thr Gly Thr 500 505 510

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Asp Glu Ile Gln Ser Lys Leu Ile Ile Ser Asn Asn Pro Trp Gly Tyr 580 585 590

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<212> DNA

<213> Pasteurella multocida

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